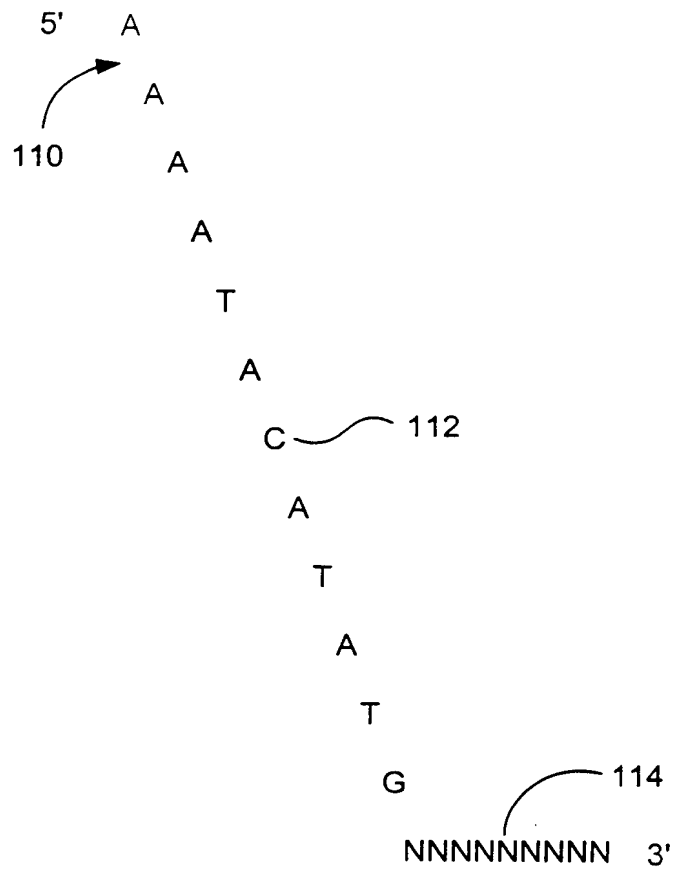


FIG. 1



CGCAATTAGAAATGCTCTCGCAAGCCGTCGTGTAGGAGACGGGTTCGGAGC

A L I F T M A F G S T S S A Q A S

100

FIG. 2A

A L I F T M A F G S T S S A Q A S
 GCGTTAATCTTTACGATGGCGTTCGGCAGCACATCCTCTGCCCAAGCCTCG

3' NNNNNNNNN

124

102

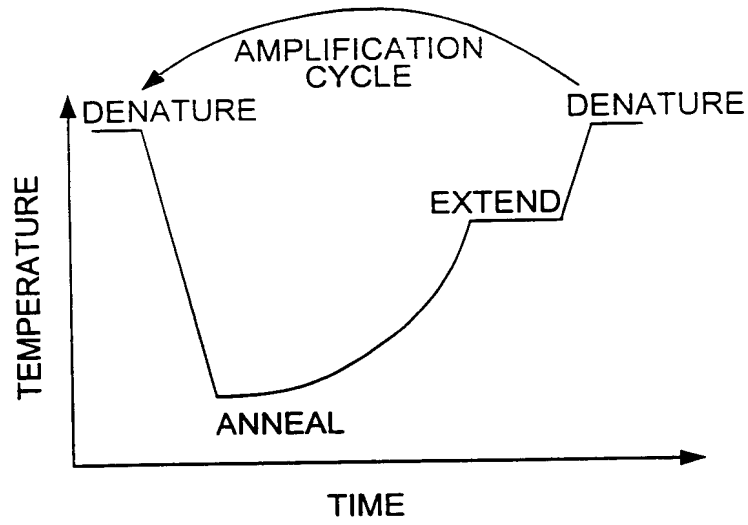
A
T
T
A
A
T
C
A
T
C
T
T
C
G
A
A
T
A 5'

122

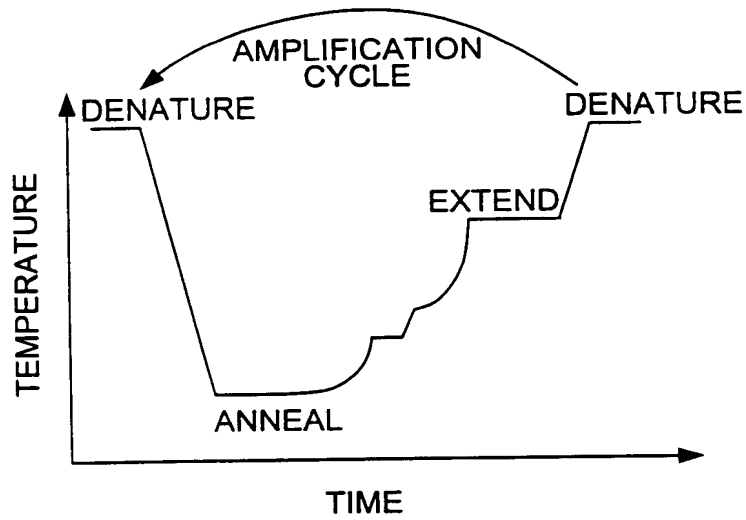
120

FIG. 2B

A



B



C

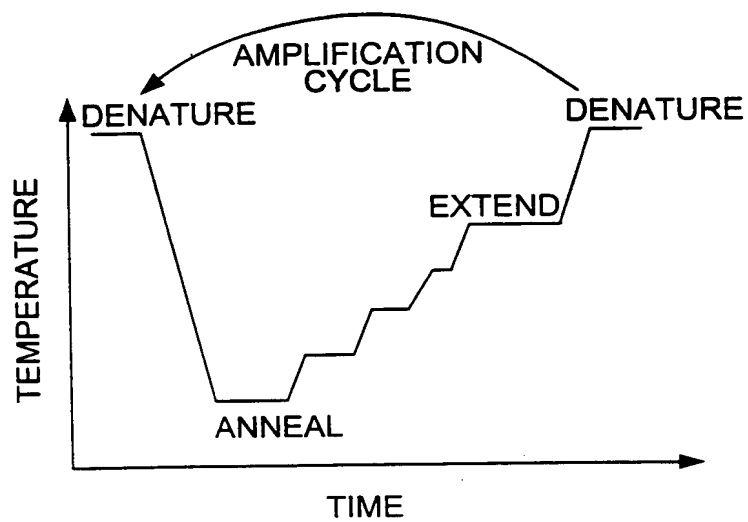


FIG. 3

1 2 3 4 5 6 7

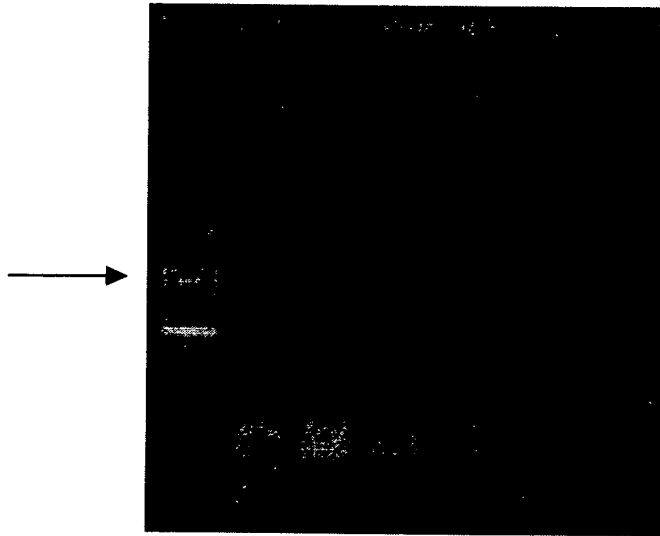


FIGURE 4

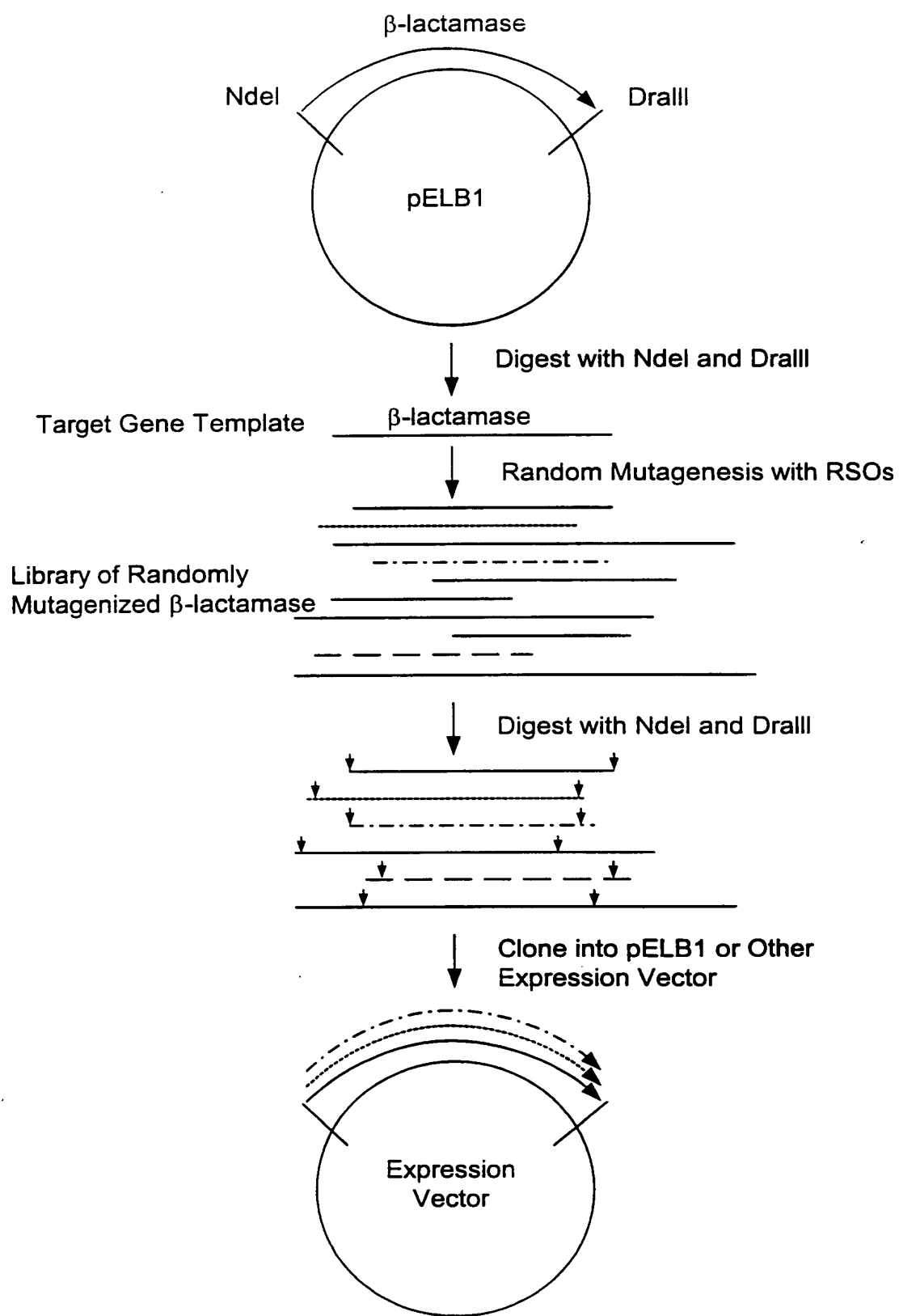


FIG. 5